

SEQUENCE LISTING

<110> Dulac, Catherine
Axel, Richard

<120> Cloning Of Vertebrate Pheromone Receptors And Uses
Thereof

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<150> 08/731,745

<151> 1996-10-18

<160> 21

<170> PatentIn Ver. 2.1

<210> 1

<211> 530

<212> DNA

<213> Homo sapiens

<400> 1

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<213> Rattus sp.

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 <213> Rattus sp.

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 Thr Asp Leu Pro Ile Gly Leu Leu Ser Leu Ile Asn Leu Leu Met Leu
 50 55 60

Leu Met Thr Ala Phe Ile Ala Thr Asp Thr Phe Ile Ser Trp Arg Gly
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 Trp Asp Asp Ile Ile Cys Lys Ser Leu Leu Tyr Leu Tyr Arg Thr Phe
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 Arg Gly Leu Ser Leu Cys Thr Ser Cys Leu Leu Ser Val Leu Gln Ala
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 Ile Ile Leu Ser Pro Arg Ser Ser Cys Leu Ala Lys Phe Lys His Lys
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 Pro Ser His His Ile Ser Cys Ala Ile Leu Ser Leu Ser Val Leu Tyr
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 Met Phe Ile Ser Ser His Leu Leu Val Ser Ile Ile Ala Thr Pro Asn
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 Leu Thr Thr Asn Asp Phe Ile His Val Thr Gln Trp Cys Ser Ile Leu
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 Pro Met Ser Tyr Leu Met Gln Ser Met Phe Ser Thr Leu Leu Ala Ile
 180 185 190
 Arg Asp Val Phe Leu Ile Ser Leu Met Val Leu Ser Thr Trp Tyr Met
 195 200 205
 Val Ala Leu Leu Cys Arg His Arg Lys Gln Thr Arg His Leu Gln Gly
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 Thr Ser Leu Ser Pro Lys Ala Ser Pro Glu Gln Arg Ala Thr Arg Ser
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 Ser Tyr Gln Leu Phe Met Val His Ile Tyr Ala Thr Val Ser Pro Phe
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<210> 9
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 Leu Leu Phe Asn Ile Phe Lys Phe Ile His Gly Gln Arg Ser Arg Leu
 35 40 45
 Thr Asp Leu Pro Ile Gly Leu Leu Ser Leu Ile Asn Leu Leu Met Leu
 50 55 60
 Leu Ile Met Ala Cys Ile Ala Thr Asp Ile Phe Ile Ser Cys Arg Arg
 65 70 75 80
 Trp Asp Asp Ile Ile Cys Lys Ser Leu Leu Tyr Leu Tyr Arg Thr Phe
 85 90 95
 Arg Gly Leu Ser Leu Ser Thr Thr Cys Leu Leu Ser Val Leu Gln Ala
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 Ile Ile Leu Ser Pro Arg Ser Ser Cys Leu Ala Lys Tyr Lys His Lys
 115 120 125
 Pro Pro His His Ile Phe Cys Ala Met Leu Phe Leu Ser Val Leu Tyr
 130 135 140
 Met Phe Ile Ser Ser His Leu Leu Leu Ser Ile Ile Ala Thr Pro Asn
 145 150 155 160
 Leu Thr Thr Asn Asp Phe Ile His Val Ser Gln Ser Cys Ser Ile Leu
 165 170 175
 Pro Met Ser Tyr Leu Met Gln Ser Met Phe Ser Thr Leu Leu Ala Ile
 180 185 190
 Arg Asn Val Phe Leu Ile Ser Leu Ile Val Leu Ser Thr Trp Tyr Met
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 Val Ala Leu Leu Cys Arg His Arg Lys Gln Thr Arg His Leu Gln Asp

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Tyr Val Arg Val Leu Asn Val
305 310

<213> Rattus sp.

Met Met Asn Lys Asn Ser Arg Val His Thr Asp Ser Thr Ile Arg Asn
1 5 10 15

Thr Phe Ser Thr Glu Ile Gly Ile Gly Ile Leu Ala Asn Ser Phe Leu
20 25 30

Leu Leu Phe His Ile Phe Lys Phe Ile Arg Gly Gln Arg Ser Asp Leu
35 40 45

Thr Asp Leu Pro Ile Gly Leu Leu Ser Leu Ile His Leu Leu Met Leu
50 55 60

Leu Met Gly Ala Phe Ile Ala Ile Asp Ile Phe Ile Ser Trp Arg Gly
65 70 75 80

Trp Asp Asp Ile Ile Cys Lys Phe Leu Val Tyr Leu Tyr Arg Ser Phe
85 90 95

Arg Gly Leu Ser Leu Cys Thr Thr Cys Met Leu Ser Val Leu Gln Ala
100 105 110

Met Phe Ser Glu Val Ser Val Gly Ile Leu Ala Asn Ser Ile Leu Phe
20 25 30

Phe Gly His Leu Cys Met Leu Leu Gly Glu Asn Lys Pro Lys Pro Ile
35 40 45

His Leu Tyr Ile Ala Ser Leu Ser Leu Thr Gln Leu Met Leu Leu Ile
50 55 60

Thr Met Gly Leu Ile Ala Ala Asp Met Phe Ile Ser Gln Gly Ile Trp
65 70 75 80

Asp Ser Thr Ser Cys Gln Ser Leu Ile Tyr Leu His Arg Leu Ser Arg
85 90 95

Gly Phe Thr Leu Ser Ala Ala Cys Leu Leu Asn Val Phe Trp Met Ile
100 105 110

Thr Leu Ser Ser Lys Lys Ser Cys Leu Thr Lys Phe Lys His Asn Ser
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Pro His His Ile Ser Gly Ala Phe Leu Leu Leu Cys Val Leu Tyr Met
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Cys Phe Ser Ser His Leu Ile Leu Ser Ile Ile Ala Thr Pro Asn Leu
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Thr Ser Asp Asn Phe Met Tyr Val Thr Lys Ser Cys Ser Phe Leu Pro
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Met Cys Tyr Ser Arg Thr Ser Met Phe Ser Thr Thr Ile Ala Val Arg
180 185 190

Glu Ala Phe Phe Ile Gly Leu Met Ala Leu Ser Ser Gly Tyr Leu Val
195 200 205

Ala Phe Leu Trp Arg His Arg Lys Gln Ala Gln His Leu His Ser Thr
210 215 220

Gly Leu Ser Ser Lys Ser Ser Pro Glu Gln Arg Ala Thr Glu Thr Ile
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Leu Leu Leu Met Ser Phe Phe Val Val Leu Tyr Ile Leu Glu Asn Val
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Val Phe Tyr Ser Ser Arg Met Phe Lys Asp Gly Ser Thr Phe Tyr Cys
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Val Gln Ile Ile Val Ser His Ser Tyr Ala Thr Val Ser Ser Phe Val
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Ala Arg Ile Ile Asn Asn
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<210> 12
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 <213> Rattus sp.

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 20 25 30

Thr Ile Gly Leu Ile Ala Ala Asp Met Phe Met Ser Arg Gly Arg Trp
 35 40 45

Asp Ser Thr Thr Cys Gln Ser Leu Ile Tyr Leu Asp Arg Leu Leu Arg
 50 55 60

Gly Phe Thr Leu Cys Ala Thr Cys Leu Leu Asn Val Leu Trp Thr Ile
 65 70 75 80

Thr Leu Ser Pro Arg Ser Ser Cys Leu Thr Thr Phe Lys His Lys Ser
 85 90 95

Pro His His Ile Ser Gly Ala Phe Leu Phe Phe Cys Val Leu Tyr Ile
 100 105 110

Ser Phe Gly Ser His Leu Phe Leu Ser Thr Ile Ala Thr Pro Asn Leu
 115 120 125

Thr Ser Asp Asn Phe Met Tyr Val Thr Lys Ser Cys Ser Phe Leu Pro
 130 135 140

Met Ser Tyr Ser Arg Thr Ser Met Phe Ser Thr Pro Met Ala Ile Arg
 145 150 155 160

Glu Ala Leu Leu Ile Gly Leu Ile Gly Leu Ser Ser Gly Tyr Met Val

165

170

175

Ala Phe Leu Trp Arg His Lys Asn Gln Ala Arg His Leu His Ser Thr
180 185 190

Ser Leu Ser Ser Lys Val Ser Pro Glu Gln Arg Ala Thr Arg Thr Ile
195 200 205

Met Ile Leu Met Ser Phe Phe Val Val Leu Tyr Ile Leu Glu Asn Val
210 215 220

Val	Phe	Tyr	Ser	Arg	Met	Thr	Phe	Lys	Asp	Gly	Ser	Met	Phe	Tyr	Cys
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Val Gln Ile Ile Val Ser His Ser Tyr Ala Thr Ile Ser Pro Phe Val
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Phe Ile Cys Thr Glu Lys Arg Ile Ile Lys Leu Trp Gly Ser Met Ser
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Ser Arg Ile Val Ser Ile
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<210> 13
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<400> 13

Met Arg Arg Ile Ser Thr Leu Tyr Gly Val Val Asp Lys Gln Ala Ile
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Phe Phe Ser Glu Val Val Ile Gly Ile Ser Phe Asn Ser Ile Leu Phe
20 25 30

Leu Phe His Ile Phe Gln Phe Leu Leu Glu Arg Arg Leu Arg Ile Thr
35 40 45

Asp Leu Ile Ile Ser Leu Leu Ala Leu Ile His Leu Gly Met Leu Thr
50 55 60

Val	Met	Gly	Phe	Arg	Ala	Val	Asp	Ile	Phe	Ala	Ser	Gln	Asn	Val	Trp
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Asn Asp Ile Lys Cys Lys Ser Leu Ala His Leu His Arg Leu Leu Arg
85 90 95

Gly	Leu	Ser	Leu	Cys	Ala	Thr	Cys	Leu	Leu	Ser	Ile	Phe	Gln	Ala	Ile	
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Thr	Leu	Ser	Pro	Arg	Ser	Ser	Cys	Leu	Ala	Lys	Phe	Lys	Tyr	Lys	Ser	
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Thr	Gln	His	Ser	Leu	Cys	Ser	Leu	Leu	Val	Leu	Trp	Ala	Phe	Tyr	Met	
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Met	Asp	Tyr	Ile	Thr	Arg	His	Leu	Phe	Phe	Ile	Leu	Gly	Ile	Phe	Arg	
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Asp	Val	Ser	Phe	Ile	Gly	Leu	Met	Ala	Leu	Ser	Ser	Gly	Tyr	Met	Val	
			195					200			205					
Ala	Leu	Leu	Cys	Arg	His	Arg	Lys	Gln	Ala	Gln	His	Leu	His	Arg	Thr	
			210					215			220					
Ser	Leu	Ser	Pro	Lys	Ala	Ser	Pro	Glu	Gln	Arg	Ala	Thr	Arg	Thr	Ile	
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<210> 14
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<212> PRT
<213> Rattus sp.
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<400> 14

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Leu Cys Met Phe Phe Glu Glu Asn Arg Ser Lys Pro Ile Asp Leu Cys
35 40 45

Ile Ala Phe Leu Ser Leu Thr Gln Leu Met Leu Leu Val Thr Met Gly
50 55 60

Leu Ile Ala Ala Asp Met Phe Met Ala Gln Gly Ile Trp Asp Ile Thr
65 70 75 80

Thr Cys Arg Ser Leu Ile Tyr Phe His Arg Leu Leu Arg Gly Phe Asn
85 90 95

Leu Cys Ala Ala Cys Leu Leu His Ile Leu Trp Thr Phe Thr Leu Ser
100 105 110

Pro Arg Ser Ser Cys Leu Thr Lys Phe Lys His Lys Ser Pro His His
115 120 125

Ile Ser Gly Ala Tyr Leu Phe Phe Cys Val Leu Tyr Met Ser Phe Ser
130 135 140

Ser His Leu Phe Val Leu Val Ile Ala Thr Ser Asn Leu Thr Ser Asp
145 150 155 160

His Phe Met Tyr Val Thr Gln Ser Cys Ser Leu Leu Pro Met Ser Tyr
165 170 175

Ser Arg Thr Ser Thr Phe Ser Leu Leu Met Val Thr Arg Glu Val Phe
180 185 190

Leu Ile Ser Leu Met Ala Leu Ser Ser Gly Tyr Met Val Thr Leu Leu
195 200 205

Trp Arg His Lys Lys Gln Ala Gln His Leu His Ser Thr Arg Leu Ser
210 215 220

Ser Lys Ala Ser Pro Gln Gln Arg Ala Thr Arg Thr Ile Leu Leu Leu
225 230 235 240

Met Thr Phe Phe Val Val Phe Tyr Ile Leu Gly Thr Val Ile Phe His
245 250 255

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150

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160

Phe Met Cys Leu Phe Pro Phe Ala Ala Leu Thr Leu Leu
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<210> 16

<211> 71

<212> PRT

<213> Rattus sp.

<400> 16

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 20 25 30

Tyr Leu Ser Gln Arg Arg Trp Glu Gln Leu Asp Pro Ser Gly Arg Leu
 35 40 45

Cys Thr Phe Phe Gly Leu Thr Met Thr Val Phe Gly Leu Ser Ser Leu
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Leu Val Ala Ser Ala Met Ala
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<210> 17

<211> 74

<212> PRT

<213> Rattus sp.

<400> 17

Gly Gln Arg Ser Arg Leu Thr Asp Leu Pro Ile Gly Leu Leu Ser Leu
 1 5 10 15

Ile Asn Leu Leu Met Leu Leu Ile Met Ala Cys Ile Ala Thr Asp Ile
 20 25 30

Phe Ile Ser Cys Arg Arg Trp Asp Asp Ile Ile Cys Lys Ser Leu Leu
 35 40 45

Tyr Leu Tyr Arg Thr Phe Arg Gly Leu Ser Leu Ser Thr Thr Cys Leu
 50 55 60

Leu Ser Val Leu Gln Ala Ile Ile Leu Ser
 65 70

1990-1991		1991-1992		1992-1993		1993-1994		1994-1995		1995-1996		1996-1997		1997-1998		1998-1999		1999-2000		2000-2001		2001-2002		2002-2003		2003-2004		2004-2005		2005-2006		2006-2007		2007-2008		2008-2009		2009-2010		2010-2011		2011-2012		2012-2013		2013-2014		2014-2015		2015-2016		2016-2017		2017-2018		2018-2019		2019-2020		2020-2021		2021-2022		2022-2023		2023-2024		2024-2025		2025-2026		2026-2027		2027-2028		2028-2029		2029-2030		2030-2031		2031-2032		2032-2033		2033-2034		2034-2035		2035-2036		2036-2037		2037-2038		2038-2039		2039-2040		2040-2041		2041-2042		2042-2043		2043-2044		2044-2045		2045-2046		2046-2047		2047-2048		2048-2049		2049-2050		2050-2051		2051-2052		2052-2053		2053-2054		2054-2055		2055-2056		2056-2057		2057-2058		2058-2059		2059-2060		2060-2061		2061-2062		2062-2063		2063-2064		2064-2065		2065-2066		2066-2067		2067-2068		2068-2069		2069-2070		2070-2071		2071-2072		2072-2073		2073-2074		2074-2075		2075-2076		2076-2077		2077-2078		2078-2079		2079-2080		2080-2081		2081-2082		2082-2083		2083-2084		2084-2085		2085-2086		2086-2087		2087-2088		2088-2089		2089-2090		2090-2091		2091-2092		2092-2093		2093-2094		2094-2095		2095-2096		2096-2097		2097-2098		2098-2099		2099-2100		2100-2101		2101-2102		2102-2103		2103-2104		2104-2105		2105-2106		2106-2107		2107-2108		2108-2109		2109-2110		2110-2111		2111-2112		2112-2113		2113-2114		2114-2115		2115-2116		2116-2117		2117-2118		2118-2119		2119-2120		2120-2121		2121-2122		2122-2123		2123-2124		2124-2125		2125-2126		2126-2127		2127-2128		2128-2129		2129-2130		2130-2131		2131-2132		2132-2133		2133-2134		2134-2135		2135-2136		2136-2137		2137-2138		2138-2139		2139-2140		2140-2141		2141-2142		2142-2143		2143-2144		2144-2145		2145-2146		2146-2147		2147-2148		2148-2149		2149-2150		2150-2151		2151-2152		2152-2153		2153-2154		2154-2155		2155-2156		2156-2157		2157-2158		2158-2159		2159-2160		2160-2161		2161-2162		2162-2163		2163-2164		2164-2165		2165-2166		2166-2167		2167-2168		2168-2169		2169-2170		2170-2171		2171-2172		2172-2173		2173-2174		2174-2175		2175-2176		2176-2177		2177-2178		2178-2179		2179-2180		2180-2181		2181-2182		2182-2183		2183-2184		2184-2185		2185-2186		2186-2187		2187-2188		2188-2189		2189-2190		2190-2191		2191-2192		2192-2193		2193-2194		2194-2195		2195-2196		2196-2197		2197-2198		2198-2199		2199-2200		2200-2201		2201-2202		2202-2203		2203-2204		2204-2205		2205-2206		2206-2207		2207-2208		2208-2209		2209-2210		2210-2211		2211-2212		2212-2213		2213-2214		2214-2215		2215-2216		2216-2217	
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Lys Cys Lys Ser Leu Ala His Leu His Arg Leu Leu Arg Gly Leu Ser
1 5 10 15

Pro Arg Ser Ser Cys Leu Ala Lys Ser Thr Gln His Ser Leu Cys Ser
35 40 45

Leu Leu Val Leu Trp Ala Phe Tyr Met Ser Cys Gly Thr His Tyr Ser
50 55 60

Phe Thr Ile Val Ala Asp Tyr Asn Phe Ser Ser Arg Ser Leu Ile Phe
65 70 75 80

Val Thr Glu Ser Cys Ile Ile Leu Pro Met Asp Tyr Ile Thr Arg Asp
85 90 95

Leu Phe Phe Ile Leu Gly Ile Phe Arg Asp Val Ser Phe Ile Gly Leu
100 105 110

Met Ala Leu Ser Ser Gly Tyr Met Val Ala Leu Leu Cys Arg His Arg
115 120 125

Lys Gly Ala Gln His Leu His Arg Thr Ser Leu Ser Pro Lys Ala Ser
130 135 140

Pro	Glu	Gln	Arg	Ala	Thr	Arg	Thr	Ile	Leu	Leu	Leu	Met	Ser	Phe	Phe
145					150					155					160

Val Leu Met Tyr Cys Leu Asp Cys Thr Ile Ser Ala Ser Arg
165 170

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<210> 19
<211> 32
<212> PRT
<213> Rattus sp.
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<222> (31)
<223> Xaa at position 31 is His or Tyr

<220>
<221> VARIANT
<222> (32)
<223> Xaa at position 32 is Lys or Asn

<400> 19
Arg Gly Xaa Xaa Leu Xaa Xaa Xaa Cys Xaa Leu Xaa Xaa Xaa Xaa Xaa
1 5 10 15

Xaa Xaa Leu Ser Xaa Xaa Xaa Ser Cys Leu Xaa Xaa Xaa Lys Xaa Xaa
20 25 30

<210> 20
<211> 21
<212> PRT
<213> Rattus sp.

<220>
<221> VARIANT
<222> (2)
<223> Xaa at position 2 is Ala or Ser or Val

<220>
<221> VARIANT
<222> (5)
<223> Xaa at position 5 is Glu or Gln

<220>
<221> VARIANT
<222> (10)
<223> Xaa at position 10 is Arg or Gln or Glu

<220>
<221> VARIANT
<222> (11)
<223> Xaa at position 11 is Thr or Ser

<220>
<221> VARIANT
<222> (13)
<223> Xaa at position 13 is Leu or Met

[illegible]

<223> Xaa at position 4 is Val or Ile or Leu

<223> Xaa at position 6 is Pro or Ser

<223> Xaa at position 7 is Phe or Leu

<223> Xaa at position 8 is Val or Leu

<223> Xaa at position 9 is Phe or Leu

1

5